

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:02:42 ; Search time 350.985
Seconds

(without alignments)
1814.018 Million cell

updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataaggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 9369276 seqs, 1007425964 residues

Total number of hits satisfying chosen parameters: 18738552

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_New:*
1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
2: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq2:*
3: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
4: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
5: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
6: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*
9: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq3:*
10: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq2:*
12: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq3:*
13: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*